

# Tomer Chen

## Bioinformatician

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### Professional Summary

Bioinformatician with expertise in RNA therapeutics, genetic engineering, and computational biology. Skilled in automation, bioinformatics pipelines, and data analysis, reducing outsourcing costs and improving efficiency. Developed Python-based RNA analysis tools and automation frameworks, cutting analysis time by 80%. Proficient in CRISPR, NGS, statistical modeling, and full-stack development, bridging biotechnology and software engineering to drive innovation.

### Skills

**Biotechnology:** PCR, WB, Transfection, Transformation, cloning, RNA design, Mars-Seq, RNA-Seq NGS data analysis, DNA/RNA/Protein extraction, CRISPR & Gene Editing, Statistical Modelling, JMP.

**Programming:** Python, C#, JavaScript, TypeScript, RESTful API Development, WPF, SQL, NoSQL (MongoDB).

**Tools:** Git, VS-Code, Visual Studio, Spyder, Azure

**Soft Skills:** Fast learner, Multitasking, Technical Writing & Documentation, Problem-Solving, Project Management, Teamwork.

### Work Experience

**Senior Scientist, Head of RNA Therapeutics** 2023-Present

#### ArtBioScience

- Developed Python-based RNA analysis software, eliminating outsourcing costs and enabling in-house analysis.
- Developed an automation framework, reducing file analysis time by up to 80%.
- Created automation software, enhancing data presentation consistency and uniformity.

**Scientist, RNA Therapeutics Department** 2021-2023

#### ArtBioScience

- Developed a Python program to optimize mRNA synthesis by removing inhibitory sequences and restriction sites, increasing yields by 20%-100% and enabling the production of previously unsynthesizable mRNAs while reducing external optimization costs.

**Scientific Consultant** 2021

#### Weizmann Institute of Science

- Advised on experimental design, data analysis, and molecular biology techniques, improving research methods.

**Postdoctoral researcher** 2019-2021

#### Weizmann Institute of Science

- Led a research team of 3 scientists focused on developing drought-resistant plants via genetic engineering.
- Designed inducible vectors and localized promoters using  $\mu$ RNA to reduce gene expression in target plant tissues.
- Generated modified Arabidopsis thaliana plants that required 30% less water than wild-type.

### Projects

#### HacherU Course

##### WPF Final Project

- Developed a gallery showcasing multiple WPF applications, leveraging OOP principles, LINQ for efficient data querying, and APIs. Utilized JSON for seamless data exchange, delivering a dynamic, responsive, user-friendly interface.

##### Full-Stack Final Project

- Designed and built a full-stack social media platform with a React-based interactive front end and a scalable .NET-powered RESTful API. Deployed the API on **Azure** and front end on **Render**, ensuring a robust, efficient architecture.

### Education

#### Courses & Certifications

<b>HackerU</b> - Dot Net Full-Stack Developer	2023-2024
<b>SoloLearn</b> - Python Developer Certification	2023
<b>SoloLearn</b> - Python Core Certification	2022

#### Academic

<b>Ph.D. in Biotechnology</b> - Weizmann Institute of Science	2014-2019
Thesis: Singlet Oxygen Synthesis Under Osmotic Shock Conditions	
<b>M.Sc. in Biotechnology</b> - Bar-Ilan University	2011-2014
Thesis: Genetic Resistance of Cucumbers Against Downy Mildew	
Professor Yehuda Halevy Prize for Exceptional Research	
<b>B.Sc. in Biotechnology, Psagot Program for Direct MSc Studies</b> - Bar-Ilan University	2010-2011
Research Project: Identification Of Resistance Genes in Tomato Using Genetic Markers	

### Publications

<i>Osmotic stress in roots drives lipoxygenase-dependent plastid remodeling through singlet oxygen production</i>	2024
<i>Lipoxygenase functions in <math>1O_2</math> production during root responses to osmotic stress</i>	2021
<i>Isolate-Dependent Inheritance of Resistance Against Pseudoperonospora cubensis in Cucumber</i>	2020
<i>Singlet Oxygen Plays an Essential Role in the Root's Response to Osmotic Stress</i>	2018

### Languages

English (Native)      Hebrew (Native)